

logo

NTDB id 615 LCA RS02545 WP 011374200.1
NTDB id 593 KW2 RS05130 WP 021037147.1
NTDB id 342514 ETT44 RS02915 WP 002988971.1
NTDB id 277 KZH43 RS10090 WP 000867601.1
NTDB id 236 SPD RS10765 WP 000867601.1
NTDB id 202 SPR RS10250 WP 000867601.1
NTDB id 167 SP RS11275 WP 000867616.1
NTDB id 507 SM12261 RS09240 WP 000867722.1
NTDB id 535 SMSK321 RS10515 WP 000867726.1
consensus



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consensus



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consensus



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consensus



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NTDB id 615 LCA RS02545 WP 011374200.1	L FVPQVRLLPVANRLSTLLPAVRIETLHANDPDQIKKITAFAQQIQLMVTITILERGVTFKKNVAVLVLGAEHMFNEA	393
NTDB id 593 KW2 RS05130 WP 021037147.1	IFVAEIDFGQDFAKNLQEKFPKEKMAFVASTTKSRKTIVEAFRKKQVSILITTSILERGVTFSSIDVFVINSEHPNFTKS	383
NTDB id 342514 ETT44 RS02915 WP 002988971.1	IFLPTIIMAESVAILKELLPAEQIACVSSQSNRKEDITAFRQGKKTILITTLERGVTFPQIDVFVLGSHHRVYSSQ	385
NTDB id 277 KZH43 RS10090 WP 000867601.1	IFASEIKKGEQLAEILQEQFPNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKS	385
NTDB id 236 SPD RS10765 WP 000867601.1	IFASEIKKGEQLAEILQEQFPNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKS	385
NTDB id 202 SPR RS10250 WP 000867601.1	IFASEIKKGEQLAEILQEQFPNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKS	385
NTDB id 167 SP RS11275 WP 000867616.1	IFASEIKKGEQLAEILQEQFPNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKS	385
NTDB id 507 SM12261 RS09240 WP 000867722.1	IFASEIKKGEQLKEITQEFPNEKIGFVSSITENRLEQVQAFRDRELITILISTTILERGVTFPCVDVFVVEANHRLFTKS	385
NTDB id 535 SMSK321 RS10515 WP 000867726.1	IFASEIKKGEQLKEILQKEFPHEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKS	385
consensus	*!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!	

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NTDB id 615 LCA RS02545 WP 011374200.1	SLIQIGRVGRSMRPTGDLFFHDGLNASIKKAIKEIQMMNKEAGL	445
NTDB id 593 KW2 RS05130 WP 021037147.1	ALIQMAGRVGRDSKRPTGLVSFFHSGKSLAMCQAQKEIKKMNRLLGGF.....	430
NTDB id 342514 ETT44 RS02915 WP 002988971.1	SLVQIAGRVGRSIDRPGDTLYFFHEGISKAMLLARKEIKEMNYKGYSHDLSTMSTN	441
NTDB id 277 KZH43 RS10090 WP 000867601.1	SLIQIGGRVGRSMRPTGDLFFHDGLNASIKKAIKEIQMMNKEAGL.....	432
NTDB id 236 SPD RS10765 WP 000867601.1	SLIQIGGRVGRSMRPTGDLFFHDGLNASIKKAIKEIQMMNKEAGL.....	432
NTDB id 202 SPR RS10250 WP 000867601.1	SLIQIGGRVGRSMRPTGDLFFHDGLNASIKKAIKEIQMMNKEAGL.....	432
NTDB id 167 SP RS11275 WP 000867616.1	SLIQIGGRVGRSMRPTGDLFFHDGLNASIKKAIKEIQMMNKEAGL.....	432
NTDB id 507 SM12261 RS09240 WP 000867722.1	SLIQIGGRVGRSMRPTGDLFFHDGLNTSIKKAIKEIQMMNKEAGL.....	432
NTDB id 535 SMSK321 RS10515 WP 000867726.1	SLIQIGGRVGRSMRPTGDLFFHDGLNTSIKKAIKEIQMMNKEAGL.....	432
consensus	*!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!	

- [X] non conserved
- [X] similar
- [X] ≥ 50% conserved